

SEQUENCE LISTING

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<110> Fletchner, J.
     Prince-Cohane, K.
     Mehta, S.
     Slusarewicz, P.
     Andjelic, S.
     Barber, B.
<120> IMPROVED HEAT SHOCK PROTEIN-BASED
 VACCINES AND IMMUNOTHERAPIES
<130> 8449-406-999
<140> 10/820,067
<141> 2004-04-08
<150> 60/462,469
<151> 2003-04-11
<150> 60/463,746
<151> 2003-04-18
<150> 60/503,417
<151> 2003-09-16
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<222> 4, 6
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<223> motif in heptamiric region recognized by heat
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Lys Thr Gly Gly Pro Ile Tyr Lys Arg
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Phe Ala Pro Gly Asn Tyr Pro Ala Leu
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Asp Pro Val Ile Asp Arg Leu Tyr Leu
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Ser Pro Gly Arg Ser Phe Ser Tyr Phe
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Tyr Pro Ala Leu Gly Leu His Glu Phe
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Thr Tyr Lys Asp Thr Val Gln Leu
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Phe Tyr Asp Gly Phe Ser Lys Val Pro Leu
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Tyr Pro His Phe Met Pro Thr Asn Leu
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Ala Pro Thr Ala Gly Ala Phe Phe
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Ser Thr Leu Pro Glu Thr Thr Val Val Arg Arg
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Phe Leu Pro Ser Asp Phe Phe Pro Ser Val
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Trp Leu Ser Leu Leu Val Pro Phe Val
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Gly Leu Ser Pro Thr Val Trp Leu Ser Val
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Asp Leu Met Gly Tyr Ile Pro Leu Val
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Leu Met Gly Tyr Ile Pro Leu Val Gly Ala
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Lys Leu Val Ala Leu Gly Ile Asn Ala Val
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Phe Leu Arg Gly Arg Ala Tyr Gly Leu
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Arg Arg Ile Tyr Asp Leu Ile Glu Leu
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Cys Leu Gly Gly Leu Leu Thr Met Val
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Ser Ser Ile Glu Phe Ala Arg Leu
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 Asp Tyr Ala Thr Leu Gly Val Gly Val
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 Leu Leu Gly Thr Leu Asn Ile Val
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 Leu Leu Met Gly Thr Leu Gly Ile Val
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 Thr Leu Gln Asp Ile Val Leu His Leu
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 Gly Leu His Cys Tyr Glu Gln Leu Val
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Ser Ala Ile Asn Asn Tyr Ala Gln Lys Leu
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His Gln Ala Ile Ser Pro Arg Thr Leu
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Gln Met Val His Gln Ala Ile Ser Pro Arg Thr Leu
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Gln Gly Ile Asn Asn Leu Asp Asn Leu
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Ser Phe Asn Cys Gly Glu Phe Phe
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Gly Arg Ala Phe Val Thr Ile Gly Lys
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Thr Glu Met Glu Lys Glu Gly Lys Ile
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Ile Leu Lys Glu Pro Val His Gly Val
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Val Glu Ala Glu Ile Ala His Gln Ile
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Ile Ile Tyr Arg Phe Leu Leu Ile
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Lys Tyr Gly Val Ser Val Gln Asp Ile
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Tyr Leu Glu Pro Gly Pro Val Thr Ala
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Leu Asn Trp Pro Arg Val Leu Trp
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Ala Val Thr Ala Ala Ile Val Trp
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Thr Pro Leu Lys Val Pro Tyr Trp
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Cys Arg Phe His Gly Asn Arg Trp
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Ser Ser Ala Ala Gly Pro Arg Trp
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Ser Leu Ile Gln Tyr Ser Arg Trp
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Thr Val Gln His Val Ala Phe Trp
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Val Gly Ser Met Glu Ser Leu Trp
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Ile Ala Thr Lys Thr Pro Lys Trp
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<211> 8
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<210> 817
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<400> 818
Thr Thr Pro Pro Asn Phe Ala Trp
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<210> 822
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<210> 823
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Ala Pro Leu Asp Arg Ile Thr Trp
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<210> 826
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<400> 826
Phe Ala Pro Leu Ile Ala His Trp
<210> 827
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Ser Trp Ile Gln Thr Phe Met Trp
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Asn Thr Trp Pro His Met Tyr Trp
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Glu Pro Leu Pro Thr Thr Leu Trp
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His Gly Pro His Leu Phe Asn Trp
<210> 831
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<400> 831
Tyr Leu Asn Ser Thr Leu Ala Trp
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Thr Leu Pro His Arg Leu Asn Trp
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Ser Ser Pro Arg Glu Val His Trp
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Asn Gln Val Asp Thr Ala Arg Trp
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His Pro Ala Ala Phe Pro Trp Trp
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Leu Leu Pro His Ser Ser Ala Trp
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Leu Glu Thr Tyr Thr Ala Ser Trp
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      "Trp" residue
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Lys Tyr Val Pro Leu Pro Pro Trp
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<210> 841
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<223> Heat shock protein binding domain with terminal
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Ala Pro Leu Ala Leu His Ala Trp
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<210> 842
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Tyr Glu Ser Leu Leu Thr Lys Trp
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<210> 843
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Ser His Ala Ala Ser Gly Thr Trp
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<400> 844
Gly Leu Ala Thr Val Lys Ser Trp
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Thr Leu Tyr Val Ser Gly Asn Trp
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His Ala Pro Phe Lys Ser Gln Trp
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Val Ala Phe Thr Arg Leu Pro Trp
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<210> 851
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Ala Ser Phe Asp Leu Leu Ile Trp
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      "Trp" residue
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Arg Met Asn Thr Glu Pro Pro Trp
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      "Trp" residue
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Lys Met Thr Pro Leu Thr Trp
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Ala Asn Ala Thr Pro Leu Leu Trp
                5
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<400> 855
Thr Ile Trp Pro Pro Pro Val Trp
               5
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Gln Thr Lys Val Met Thr Thr Trp
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<223> Heat shock protein binding domain with terminal
      "Trp" residue
<400> 857
Asn His Ala Val Phe Ala Ser Trp
                 5
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Leu His Ala Ala Xaa Thr Ser Trp
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Thr Trp Gln Pro Tyr Phe His Trp
                 5
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      "Trp" residue
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Ala Pro Leu Ala Leu His Ala Trp
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<210> 861
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      "Trp" residue
<400> 861
Thr Ala His Asp Leu Thr Val Trp
                 5
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<210> 862
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<400> 862
Asn Met Thr Asn Met Leu Thr Trp
                 5
<210> 863
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     "Trp" residue
<400> 863
Gly Ser Gly Leu Ser Gln Asp Trp
<210> 864
<211> 8
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     "Trp" residue
<400> 864
Thr Pro Ile Lys Thr Ile Tyr Trp
<210> 865
<211> 8
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      "Trp" residue
<400> 865
Ser His Leu Tyr Arg Ser Ser Trp
                 5
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<223> Heat shock protein binding domain with terminal
      "Trp" residue
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His Gly Gln Ala Trp Gln Phe Trp
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<210> 867
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Asn Leu Leu Arg Leu Thr Gly Trp
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<210> 868
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Ser Ile Ile Asn Phe Glu Lys Leu
<210> 869
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<223> Heat shock-protein binding motif to form hybrid antigen
<400> 869
His Trp Asp Phe Ala Trp Pro Trp
<210> 870
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<223> Heat shock-protein binding motif to form hybrid antigen
<400> 870
Asn Leu Leu Arg Leu Thr Gly Trp
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Phe Tyr Gln Leu Ala Leu Thr Trp
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Arg Lys Leu Phe Phe Asn Leu Arg Trp
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<210> 873
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<400> 873
Ala Leu Phe Asp Ile Glu Ser Lys Val
<210> 874
<211> 9
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<400> 874
Ile Met Asp Gln Val Pro Phe Ser Val
<210> 875
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<223> Heat shock-protein binding motif to form hybrid antigen
<400> 875
Tyr Met Asp Gly Thr Met Ser Gln Val
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<210> 876
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Thr Leu Gly Ile Val Cys Pro Ile
<210> 877
<211> 10
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Tyr Met Leu Asp Leu Gln Pro Glu Thr Thr
                 5
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Ala Leu Phe Asp Ile Glu Ser Lys Val Gly Ser Gly His Trp Asp Phe
Ala Trp Pro Trp
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<210> 879
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Arg Gly Tyr Val Tyr Gln Gly Leu
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<210> 880
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Glu Lys Leu
<210> 881
<211> 20
<212> PRT
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<223> Hybrid antigen
<400> 881
Asn Leu Leu Arg Leu Thr Gly Trp Phe Phe Arg Lys Ser Ile Ile Asn
                                     10
Phe Glu Lys Leu
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<210> 882
<211> 18
<212> PRT
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<400> 882
Asn Leu Leu Arg Leu Thr Gly Trp Arg Lys Ser Ile Ile Asn Phe Glu
                                     10
Lys Leu
<210> 883
<211> 19
<212> PRT
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Asn Leu Leu Arg Leu Thr Gly Trp Gly Ser Gly Arg Gly Tyr Val Tyr
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Gln Gly Leu
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<210> 884
<211> 20
<212> PRT
<213> Artificial Sequence
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<223> Hybrid antigen
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Tyr Gln Gly Leu
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<210> 885
<211> 18
<212> PRT
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<223> Hybrid antigen
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Asn Leu Leu Arg Leu Thr Gly Trp Arg Lys Arg Gly Tyr Val Tyr Gln
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Gly Leu
<210> 886
<211> 20
<212> PRT
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Asn Leu Leu Arg Leu Thr Gly Trp Ala Lys Val Leu Ser Ile Ile Asn
                                     10
Phe Glu Lys Leu
            20
<210> 887
<211> 19
<212> PRT
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<223> Hybrid antigen
<400> 887
Asn Leu Leu Arg Leu Thr Gly Trp Gln Leu Lys Ser Ile Ile Asn Phe
Glu Lys Leu
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<210> 888

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Asn Leu Leu Arg Leu Thr Gly Trp Phe Arg Ser Ile Ile Asn Phe Glu
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Lys Leu
<210> 889
<211> 21
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<400> 889
Asn Leu Leu Arg Leu Thr Gly Trp Phe Phe Arg Lys Ile Met Asp Gln
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Val Pro Phe Ser Val
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<210> 890
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<212> PRT
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<223> Hybrid antigen
<400> 890
Asn Leu Leu Arg Leu Thr Gly Trp Phe Phe Arg Lys Tyr Met Asp Gly
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Thr Met Ser Gln Val
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<223> Heat shock-protein binding motif to form hybrid antigen
<400> 891
Phe Ala Pro Gly Asn Tyr Pro Ala Leu
<210> 892
<211> 21
<212> PRT
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<213> Artificial Sequence
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<223> Hybrid antigen
<400> 892
Asn Leu Leu Arg Leu Thr Gly Trp Phe Phe Arg Lys Phe Ala Pro Gly
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Asn Tyr Pro Ala Leu
            20
<210> 893
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<400> 893
Asn Leu Leu Arg Leu Thr Gly Trp Phe Phe Arg Lys Glu Leu Ala Gly
                                     10
Ile Gly Ile Leu Thr Val
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<210> 894
<211> 21
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Asn Leu Leu Arg Leu Thr Gly Trp Phe Phe Arg Lys Ser Leu Leu Met
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Trp Ile Thr Gln Val
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<210> 895
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Asn Leu Leu Arg Leu Thr Gly Trp Phe Phe Arg Lys Ser Val Tyr Asp
Phe Phe Val Trp Leu
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<210> 896
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<212> PRT
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<220>
<223> Hybrid antigen
<400> 896
Gly Leu Tyr Asp Gly Met Glu His Leu Gly Ser Gly Asn Leu Leu Arg
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Leu Thr Gly Trp
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<210> 897
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<400> 897
Tyr Leu Glu Pro Gly Pro Val Thr Val Gly Ser Gly Asn Leu Leu Arg
                                     10
Leu Thr Gly Trp
            20
<210> 898
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Lys Ala Ser Glu Lys Ile Phe Tyr Val Gly Ser Gly Asn Leu Leu Arg
Leu Thr Gly Trp
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<210> 899
<211> 20
<212> PRT
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Asn Leu Leu Arg Leu Thr Gly Trp Phe Phe Arg Lys Ser Ser Trp Asp
Phe Ile Thr Val
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<210> 900
<211> 31
<212> PRT
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<220>
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<400> 900
Asn Leu Leu Arg Leu Thr Gly Trp Phe Phe Arg Lys Ser Ile Ile Asn
                5
Phe Glu Lys Leu Phe Phe Arg Lys Arg Gly Tyr Val Tyr Gly Leu
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<210> 901
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Asn Leu Leu Arg Leu Thr Gly Trp Phe Phe Arg Lys Arg Gly Tyr Val
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Tyr Gln Gly Leu Phe Phe Arg Lys Ser Ile Ile Asn Phe Glu Lys Leu
                                 25
            20
<210> 902
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Asn Leu Leu Arg Leu Thr Gly Trp Phe Phe Arg Lys Ser Ile Ile Asn
                                    10
Phe Glu Lys Leu Phe Phe Arg Lys Arg Gly Tyr Val Tyr Gln Gly Leu
                                 25
<210> 903
<211> 32
<212> PRT
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<220>
<223> Hybrid antigen
<400> 903
Asn Leu Leu Arg Leu Thr Gly Trp Phe Phe Arg Lys Arg Gly Tyr Val
                                     10
Tyr Gln Gly Leu Phe Phe Arg Lys Ser Ile Ile Asn Phe Glu Lys Leu
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<210> 904
<211> 8
<212> PRT
<213> Artificial Sequence
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<220>

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<223> Heat shock-protein binding motif to form hybrid antigen
<400> 904
Ile Ala Tyr Phe Tyr Pro Glu Leu
<210> 905
<211> 32
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<223> Hybrid antigen
<400> 905
Asn Leu Leu Arg Leu Thr Gly Trp Phe Phe Arg Lys Ser Ile Ile Asn
                5
                                     10
Phe Glu Lys Leu Phe Phe Arg Lys Arg Gly Tyr Val Tyr Gln Gly Leu
                                25
<210> 906
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<223> Heat shock-protein binding motif to form hybrid antigen
<400> 906
Arg Thr Phe Ser Phe Gln Leu Ile
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<210> 907
<211> 20
<212> PRT
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<223> Hybrid antigen
<400> 907
Asn Leu Leu Arg Leu Thr Gly Trp Phe Phe Arg Lys Arg Thr Phe Ser
Phe Gln Leu Ile
            20
<210> 908
<211> 16
<212> PRT
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<223> Hybrid antigen
<400> 908
Thr Glu Trp Thr Ser Ser Asn Val Met Glu Glu Arg Lys Ile Lys Val
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<210> 909
<211> 28
<212> PRT
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<223> Hybrid antigen
<400> 909
Asn Leu Leu Arg Leu Thr Gly Trp Phe Phe Arg Lys Thr Glu Trp Thr
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Ser Ser Asn Val Met Glu Glu Arg Lys Ile Lys Val
<210> 910
<211> 20
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<400> 910
Asn Leu Leu Arg Leu Thr Gly Trp Phe Phe Arg Lys Asp Ala Pro Ile
Tyr Thr Asn Val
<210> 911
<211> 20
<212> PRT
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<223> Hybrid antigen
<400> 911
Asn Leu Leu Arg Leu Thr Gly Trp Phe Phe Arg Lys Ser Ser Trp Asp
Phe Ile Thr Val
<210> 912
<211> 20
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<220>
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Asn Leu Leu Arg Leu Thr Gly Trp Phe Phe Arg Lys Arg Thr Phe Ser
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Phe Gln Leu Ile

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<210> 913
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<400> 913
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Tyr Pro Glu Leu
<210> 914
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<212> PRT
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<400> 914
Ser Ser Trp Asp Phe Ile Thr Val
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<210> 915
<211> 8
<212> PRT
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<223> Heat shock-protein binding motif to form hybrid antigen
<400> 915
Asp Ala Pro Ile Tyr Thr Asn Val
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<210> 916
<211> 19
<212> PRT
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<400> 916
Asn Asn Phe Thr Val Ser Phe Trp Leu Arg Val Pro Lys Val Ser Ala
Ser His Leu
<210> 917
<211> 31
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<212> PRT
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<220>
<223> Hybrid antigen
<400> 917
Asn Leu Leu Arg Leu Thr Gly Trp Phe Phe Arg Lys Asn Asn Phe Thr
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Val Ser Phe Trp Leu Arg Val Pro Lys Val Ser Ala Ser His Leu
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<210> 918
<211> 13
<212> PRT
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<223> Hybrid antigen
<400> 918
Thr Pro Pro Ala Tyr Arg Pro Pro Asn Ala Pro Ile Leu
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<210> 919
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<212> PRT
<213> Artificial Sequence
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<223> Hybrid antigen
<400> 919
His Trp Asp Phe Ala Trp Pro Trp Asn Gly Ser Gly Asn Asn Phe Thr
                5
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Val Ser Phe Trp Leu Arg Val Pro Lys Val Ser Ala Ser His Leu
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<210> 920
<211> 9
<212> PRT
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<223> Heat shock-protein binding motif to form hybrid antigen
<400> 920
Ser Val Tyr Asp Phe Phe Val Trp Leu
<210> 921
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<223> Heat shock-protein binding motif to form hybrid antigen
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<400> 921
Val Ile Tyr Gln Tyr Met Asp Asp Leu
<210> 922
<211> 21
<212> PRT
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<223> Hybrid antigen
<400> 922
Asn Leu Leu Arg Leu Thr Gly Trp Phe Phe Arg Lys Ile Leu Lys Glu
Pro Val His Gly Val
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<210> 923
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Asn Leu Leu Arg Leu Thr Gly Trp Phe Phe Arg Lys Val Ile Tyr Gln
                                     10
Tyr Met Asp Asp Leu
            20
<210> 924
<211> 21
<212> PRT
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<400> 924
Asn Leu Leu Arg Leu Thr Gly Trp Phe Phe Arg Lys Ser Leu Tyr Asn
Thr Val Ala Thr Leu
            20
<210> 925
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<212> PRT
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Asn Leu Leu Arg Leu Thr Gly Trp Phe Phe Arg Lys Thr Pro Pro Ala
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Tyr Arg Pro Pro Asn Ala Pro Ile Leu
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<210> 926
<211> 30
<212> PRT
<213> Artificial Sequence

<220>
<223> Hybrid antigen

<400> 926
Asn Asn Phe Thr Val Ser Phe Trp Leu Arg Val Pro Lys Val Ser Ala
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Ser His Leu Gly Ser Gly Asn Leu Leu Arg Leu Thr Gly Trp